

ERRATUM

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Erratum: Increased spread and replication efficiency of *Listeria monocytogenes* in organotypic brain-slices is related to multilocus variable number of tandem repeat analysis (MLVA) complex

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Erratum

The original version of this article unfortunately contained a mistake. Figures two, three and four (Figs. 1, 2 and 3 here, respectively) and their associated legends were interchanged in the HTML and PDF versions of this manuscript. The correct versions are given below. In addition, Figure Five (Fig. 4 here) was missing in the HTML version of this manuscript. The correct figure Five (Fig. 4 here) is also included below.

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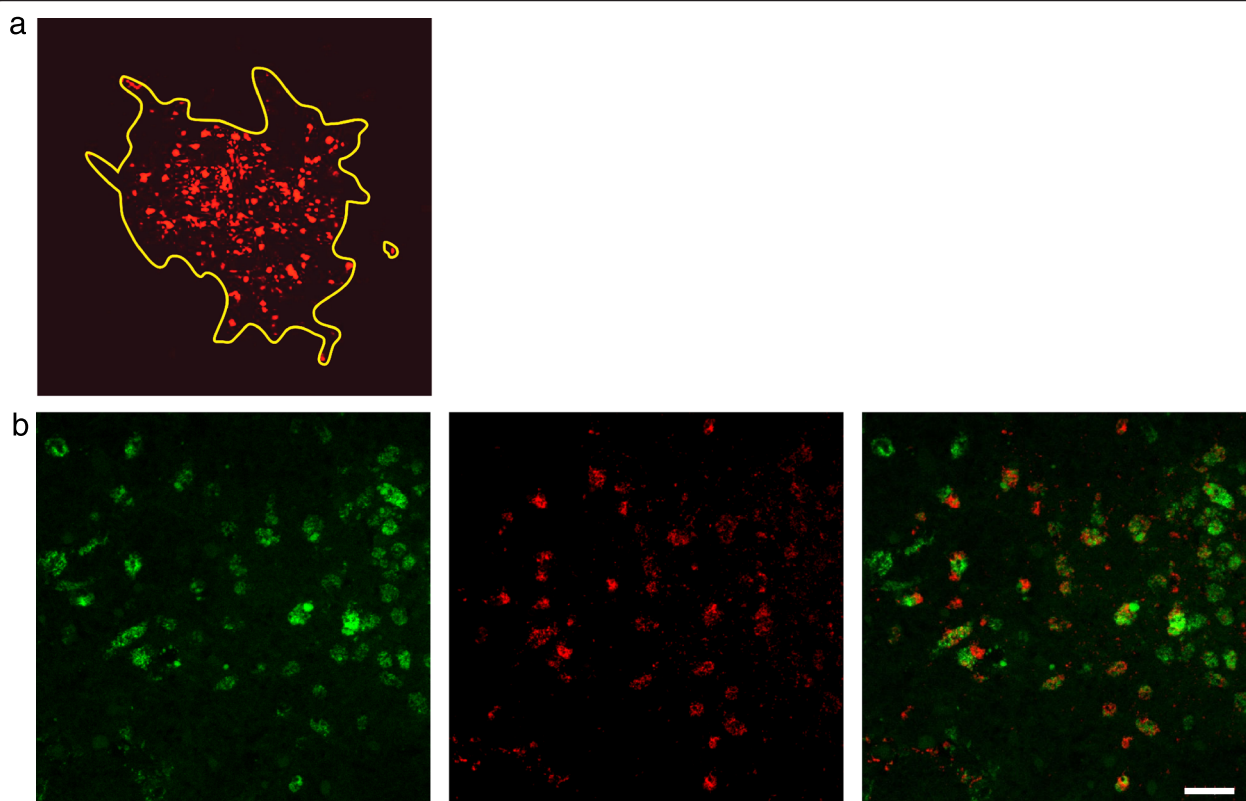
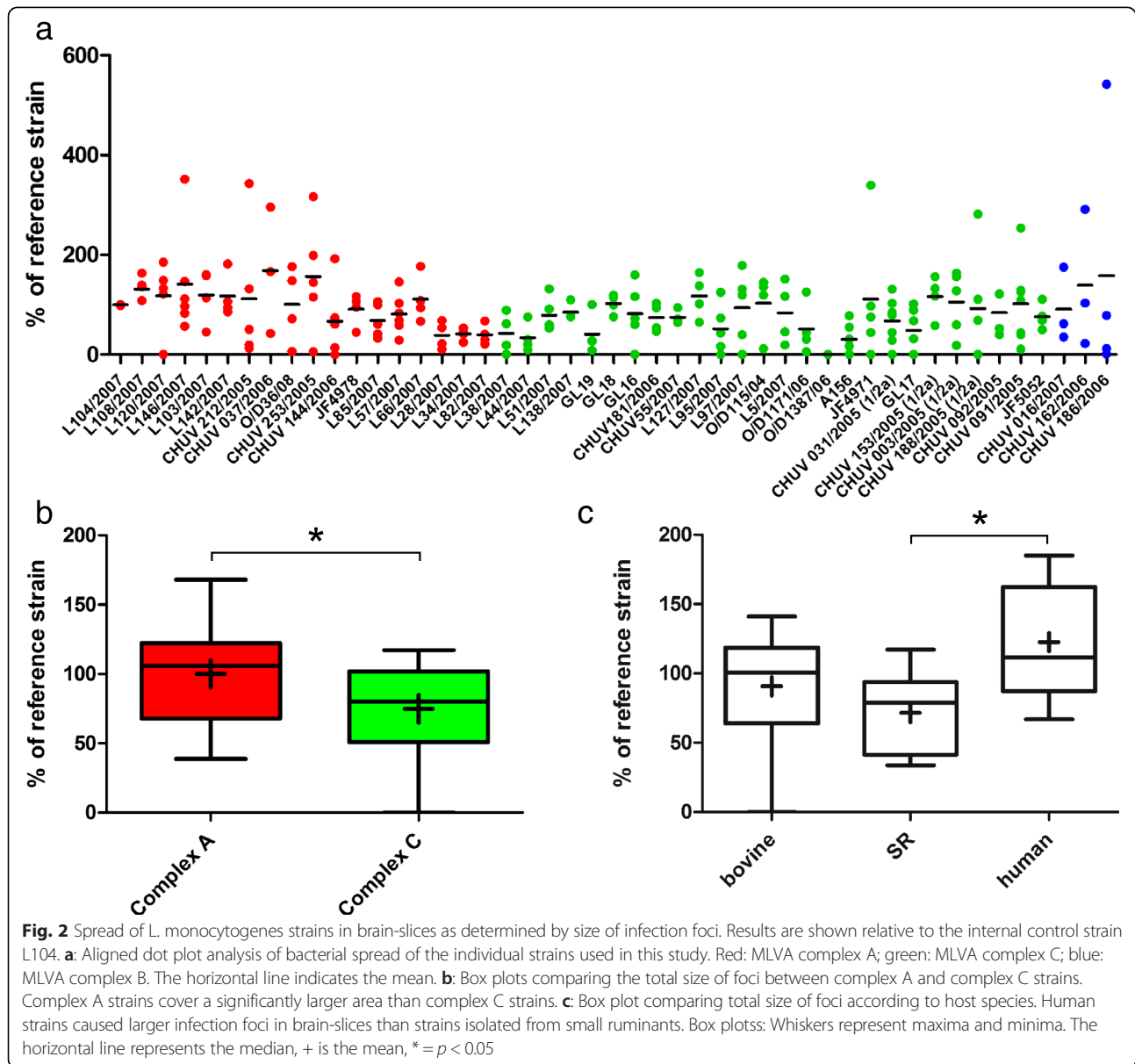


Fig. 1 Immunofluorescence stained confocal images of bacteria in infected brain-slices. **a:** Delineation of an infection focus (yellow line). *L. monocytogenes* are stained in red. The surface area covered by *L. monocytogenes* was drawn and calculated using the Fluoview software (Olympus FV10-ASW Version 03.01.01.09) Magnification 20x. **b:** Representative double-immunofluorescence of a *L. monocytogenes* infected brain-slice. The vast majority of bacteria are found within microglia. Left: Microglia are stained with CD68 in green. Center: *L. monocytogenes* in red. Right: Overlay (bar = 40 μ m)



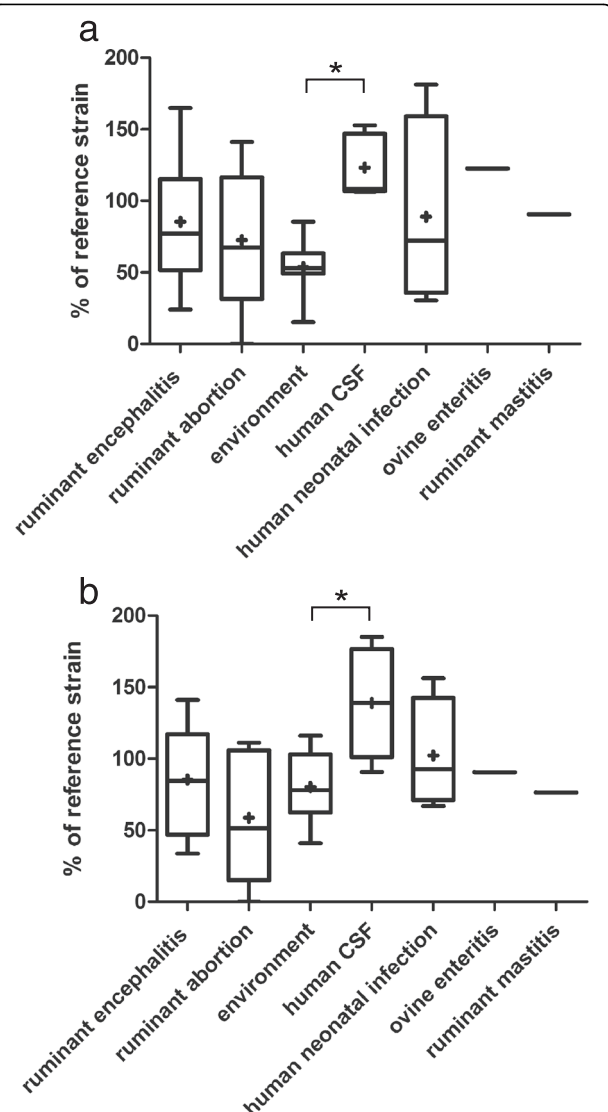


Fig. 3 CFU counts (**a**) and size of infection foci (**b**) in organotypic brain-slices infected with *L. monocytogenes* strains. Results are mapped according to the source and associated clinical infection, respectively. Data are presented as box plots, Whiskers represent maxima and minima. The vertical line represents the median, + is the mean. * = $p < 0.05$. CFU counts (**a**) and surface of bacterial spread (**b**) are shown relative to the internal control strain L104

